



SEQUENCE LISTING

<110> FOWLER, Dana M.
Broach, Jim
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Klein, Christine
Murphy, Andrew J..
Paul, Jeremy
Trueheart, Joshua

<120> YEAST CELLS ENGINEERED TO PRODUCE PHERMONE SYSTEM
PROTEIN SURROGATES, AND USES THEREFOR

<130> CPI-012CP4BCN

<140>

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<150> 08/322,137

<151> 1994-10-13

<150> 08/309,313

<151> 1994-09-20

<150> 08/190,328

<151> 1994-01-31

<150> 08/041,431

<151> 1993-03-31

<160> 133

<170> PatentIn Ver. 2.0

<210> 1

<211> 89

<212> PRT

<213> Saccharomyces cerevisiae

<400> 1

Met Arg Phe Pro Ser Ile Phe Thr Ala Val Leu Phe Ala Ala Ser Ser
1 5 10 15

Ala Leu Ala Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gln
20 25 30

Ile Pro Ala Glu Ala Val Ile Gly Tyr Leu Asp Leu Glu Gly Asp Phe
35 40 45

Asp Val Ala Val Leu Pro Phe Ser Asn Ser Thr Asn Asn Gly Leu Leu
50 55 60

Phe Ile Asn Thr Thr Ile Ala Ser Ile Ala Ala Lys Glu Glu Gly Val
65 70 75 80

Ser Leu Asp Lys Arg Glu Ala Glu Ala
85

<210> 2
<211> 76
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 2
Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr Lys Arg Glu
1 5 10 15
Ala Glu Ala Glu Ala Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro
20 25 30
Met Tyr Lys Arg Glu Ala Asp Ala Glu Ala Trp His Trp Leu Gln Leu
35 40 45
Lys Pro Gly Gln Pro Met Tyr Lys Arg Glu Ala Asp Ala Glu Ala Trp
50 55 60
His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
65 70 75

<210> 3
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aagcttaaaa gaatg

15

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aaa gaa gaa ggg gta tct ttg ctt aagctcgaga tct
Lys Glu Glu Gly Val Ser Leu Leu
1 5

37

<210> 5

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<400> 5
Lys Glu Glu Gly Val Ser Leu Leu
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<210> 6
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<222> (29)..(66)
<223> Any occurrences of n may be any nucleotide

<220>
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<400> 6
cgtgaagcctt aagcgtgagg cagaagctnn knknknknknkn nnknknknknkn nkknknknknkn 60
knknknkgtga tcattccg 77

<210> 7
<211> 19
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<221> UNSURE
<222> (7)..(19)
<223> Any occurrences of Xaa may be any amino acid

<220>
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Lys Arg Glu Ala Glu Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15
Xaa Xaa Xaa

<210> 8
<211> 36
<212> PRT
<213> Saccharomyces cerevisiae

<400> 8

Met Gln Pro Ser Thr Ala Thr Ala Ala Pro Lys Glu Lys Thr Ser Ser
1 5 10 15
Glu Lys Lys Asp Asn Tyr Ile Ile Lys Gly Val Phe Trp Asp Pro Ala
20 25 30
Cys Val Ile Ala
35

<210> 9

<211> 19

<212> DNA

<213> Artificial Sequence

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sequence

<400> 9

aagctttcga atagaaatg

19

<210> 10

<211> 36

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<400> 10

gcc gct cca aaa gaa aag acc tcg agc tcgcttaag

36

Ala Ala Pro Lys Glu Lys Thr Ser Ser

1

5

<210> 11

<211> 9

<212> PRT

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<400> 11

Ala Ala Pro Lys Glu Lys Thr Ser Ser

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5

<210> 12

<211> 79

<212> DNA

<213> Artificial Sequence

D.
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<400> 12

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gtgttattgc ttaagtacg

79

<210> 13

<211> 22

<212> PRT

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Ser Ser Glu Lys Lys Asp Asn Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15Xaa Xaa Cys Val Ile Ala
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<210> 14

<211> 34

<212> DNA

<213> Artificial Sequence

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<400> 14

gttaagaacc atatactagt atcaaaaatg tctg

34

<210> 15

<211> 35

<212> DNA

<213> Artificial Sequence

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D1
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sequence

<400> 15
tgatcaaaat ttactagttt gaaaaagtaa ttctg

35

<210> 16
<211> 28
<212> DNA
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sequence

<400> 16
ggcaaaatac tagtaaaatt ttcatgtc

28

<210> 17
<211> 34
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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 17
ggcccttaac aactagtggt cgcattatat ttac

34

<210> 18
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<212> DNA
<213> Artificial Sequence

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ctaaagaaga aggggtatct ttgcttaagc tcgagatctc gactgataac aacagtgtag 60

<210> 19
<211> 31
<212> DNA
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<220>
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sequence

<400> 19
catacacaat ataaagcttt aaaagaatga g

31

D-
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<211> 25
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gctacttaag cgtgaggcag aagct

25

<210> 21
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sequence

<400> 21
cggatgatca

10

<210> 22
<211> 41
<212> DNA
<213> Artificial Sequence

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sequence

<400> 22
ccaaaataag tacaaagctt tcgaatagaa atgcaaccat c

41

<210> 23
<211> 59
<212> DNA
<213> Artificial Sequence

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sequence

<400> 23
gccgctccaa aagaaaagac ctcgagctcg cttaagttct gcgtacaaaa acgttggtc

59

<210> 24
<211> 26
<212> DNA
<213> Artificial Sequence

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<400> 24

ggtactcgcgag tgaaaagaag gacaac

26

<210> 25

<211> 20

<212> DNA

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 25

cgacttaag caataacaca

20

<210> 26

<211> 28

<212> DNA

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 26

cgatgaagctt aagcgtgagg cagaagct

28

<210> 27

<211> 57

<212> DNA

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<222> (1)..(57)

<223> Any occurrences of N may be any nucleotide

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 27

cgatgatca mnnnnnnnnnnn nnnnnnnnnnnn nnnnnnnnnnnn mnnnnnnnnna gcttctg

57

<210> 28

<211> 26

<212> DNA

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<400> 28

ggtactcgag tgaaaagaag gacaac

26

<210> 29

<211> 60

<212> DNA

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<223> Any occurrences of N may be any nucleotide

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<400> 29

cgtacttaag caataacaca mnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn mnnngttgtcc 60

<210> 30

<211> 34

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 30

gggaagctta tgccgagatc gtgctgccag ccgc

34

<210> 31

<211> 32

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 31

ggggaagact tctgccctgc gccgctgctg cc

32

<210> 32

<211> 36

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 32

ggggaagacc cgcaggaggc agaagcttgg ttgcag

36

<210> 33

<211> 27

<212> DNA

<213> Artificial Sequence

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sequence

<400> 33

gggagatctt cagtacattg gttggcc

27

<210> 34

<211> 32

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 34

Arg Asn Ser Ser Ser Ser Gly Ser Ser Gly Ala Gly Gln Lys Arg Glu
1 5 10 15Ala Glu Ala Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
20 25 30

<210> 35

<211> 29

<212> DNA

<213> Homo sapiens

<400> 35

ccgcgtctca catgcccaag aagaagccg

29

<210> 36

<211> 24

<212> DNA

<213> Homo sapiens

<400> 36

ccgtctagat gctggcagcg tggg

24

<210> 37

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
sequence

<400> 37

ttaagcgtga ggcagaagct tatcgata

28

<210> 38

<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<400> 38

cgactccgt cttcgaatag ctatctag

28

<210> 39

<211> 71

<212> DNA

<213> Artificial Sequence

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<220>

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<400> 39

ctggatgcga agacagctnn knknknknkn nnknknknkn nkknknknkn knnktgatca 60

gtctgtgacg c

71

<210> 40

<211> 17

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 40
gcgtcacaga ctgatca

17

<210> 41
<211> 56
<212> DNA
<213> Artificial Sequence

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sequence

<400> 41
gccgtcagta aagcttggca ttggttgcag cctatgtact gatcagtcctg tgacgc 56

<210> 42
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>
<221> CDS
<222> (1)..(39)

<400> 42
tgg cat tgg ttg cag cta aaa cct ggc caa cca atg tac 39
Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
1 5 10

<210> 43
<211> 13
<212> PRT
<213> Saccharomyces cerevisiae

<400> 43
Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
1 5 10

<210> 44
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
sequence

<400> 44
ctggatgcga agactcagct 20

<210> 45

81
cont.

<211> 69
<212> DNA
<213> Artificial Sequence

<220>
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<400> 45
cggatgatca gtacattggt tggcaggtt ttagctgcaa ccaatgccaa gctgagtctt 60
cgcatccag 69

<210> 46
<211> 39
<212> DNA
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<221> CDS
<222> (1)..(39)

<400> 46
tgg cat tgg cta cag cta acg cct ggg caa cca atg tac 39
Trp His Trp Leu Gln Leu Thr Pro Gly Gln Pro Met Tyr
1 5 10

<210> 47
<211> 13
<212> PRT
<213> Saccharomyces cerevisiae

<400> 47
Trp His Trp Leu Gln Leu Thr Pro Gly Gln Pro Met Tyr
1 5 10

<210> 48
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>
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<222> (1)..(39)

<400> 48
tgg cat tgg ctg gag ctt atg cct ggc caa cca tta tac 39
Trp His Trp Leu Glu Leu Met Pro Gly Gln Pro Leu Tyr
1 5 10

<210> 49
<211> 13
<212> PRT

<213> Saccharomyces cerevisiae

<400> 49

Trp His Trp Leu Glu Leu Met Pro Gly Gln Pro Leu Tyr
1 5 10

<210> 50

<211> 39

<212> DNA

<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 50

tgg cat tgg atg gag cta aga cct ggc caa cca atg tac
Trp His Trp Met Glu Leu Arg Pro Gly Gln Pro Met Tyr
1 5 10

39

<210> 51

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 51

Trp His Trp Met Glu Leu Arg Pro Gly Gln Pro Met Tyr
1 5 10

<210> 52

<211> 33

<212> DNA

<213> Artificial Sequence

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<221> CDS

<222> (1)..(33)

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 52

tat gct ctg ttt gtt cat ttt ttt gat att ccg
Tyr Ala Leu Phe Val His Phe Phe Asp Ile Pro
1 5 10

33

<210> 53

<211> 11

<212> PRT

<213> Artificial Sequence

<400> 53

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Tyr Ala Leu Phe Val His Phe Phe Asp Ile Pro
1 5 10

<210> 54

<211> 33

<212> DNA

<213> Artificial Sequence

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<221> CDS

<222> (1)..(33)

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 54

ttt aag ggt cag gtg cgt ttt gtg gtt ctt gct
Phe Lys Gly Gln Val Arg Phe Val Val Leu Ala
1 5 10

33

<210> 55

<211> 11

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 55

Phe Lys Gly Gln Val Arg Phe Val Val Leu Ala
1 5 10

<210> 56

<211> 33

<212> DNA

<213> Artificial Sequence

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<221> CDS

<222> (1)..(33)

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<400> 56

ctt atg tct ccg tct ttt ttt ttt ttg cct gcg
Leu Met Ser Pro Ser Phe Phe Phe Leu Pro Ala
1 5 10

33

<210> 57

<211> 11
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 57

Leu Met Ser Pro Ser Phe Phe Phe Leu Pro Ala
1 5 10

<210> 58

<211> 27

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 58

cgggatccga tgcaattttc aacatgc

27

<210> 59

<211> 23

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 59

gctctagatg ctactgatcc cgc

23

<210> 60

<211> 18

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 60

cgccgcatga ctccattg

18

<210> 61

<211> 26

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 61

ggggtagcaa taggttcttt ctagg

26

<210> 62

<211> 35

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 62
ggtgggaggg tgctctctag aaggaagtgt tcacc 35

<210> 63
<211> 41
<212> DNA
<213> Artificial Sequence

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<400> 63
gcccaggaga ccagaccatg gactccttca attataccac c 41

<210> 64
<211> 42
<212> DNA
<213> Artificial Sequence

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<400> 64
ccccttaagc gtgaggcaga agctactctg caaaagaaga tc 42

<210> 65
<211> 29
<212> DNA
<213> Artificial Sequence

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<400> 65
gaagatcttc agcggccgag ttgcatgtc 29

<210> 66
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
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sequence

<400> 66
gatatattaa ggtaggaaac catgggggtgt acagtgag 38

<210> 67
<211> 34
<212> DNA
<213> Saccharomyces cerevisiae

<400> 67
cgagcgctcg agggaacgta taattaaagt agtg

34

<210> 68
<211> 34
<212> DNA
<213> Saccharomyces cerevisiae

<400> 68
gcgcggtacc aagcttcaat tcgagataat accc

34

<210> 69
<211> 24
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<400> 69
cccgaatcca ccaatttctt tacg

24

<210> 70
<211> 27
<212> DNA
<213> Artificial Sequence

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sequence

<400> 70
gcggcgctcga cgcggccgcg taacagt

27

<210> 71
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
sequence

<400> 71
ctgctggagc tccgcctgct gctgctgggt gctggag

37

Gen.

<210> 72
<211> 43
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 72
ctgctggtcg acgcggccgc gggggttcct tcttagaagc agc

43

<210> 73
<211> 30
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 73
gggctcgagc cttcttagag cagctcgtag

30

<210> 74
<211> 37
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 74
ctgctggagc tcaagttgct gctgttgggt gctgggg

37

<210> 75
<211> 44
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 75
ctgctggtcg acgcggccgc gcccctcaga agaggccgcg gtcc

44

<210> 76
<211> 29
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic sequence

<400> 76

gggctcgagc ctcagaagag gdcgcagtc

29

<210> 77

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic sequence

<400> 77

ctgctggagc tcaagctgct gctactcggg gctggag

37

<210> 78

<211> 49

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic sequence

<400> 78

ctgctggtcg acgcggccgc cactaacatc catgcttctc aataaagtc

49

<210> 79

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic sequence

<400> 79

gggctcgagc atgcttctca ataaagtcca c

31

<210> 80

<211> 19

<212> DNA

<213> Artificial Sequence

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<400> 80

D.
cont.

gcatccatca ataattcag

19

<210> 81

<211> 23

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<400> 81

gaaacaatgg atccacttct tac

23

<210> 82

<211> 66

<212> PRT

<213> Saccharomyces cerevisiae

<400> 82

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro
1 5 10 15Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Leu Gly
35 40 45Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu
50 55 60His Gln
65

<210> 83

<211> 65

<212> PRT

<213> Saccharomyces cerevisiae

<400> 83

Met Gly Cys Leu Gly Thr Ser Lys Thr Glu Asp Gln Arg Asn Glu Glu
1 5 10 15Lys Ala Gln Arg Glu Ala Asn Lys Lys Ile Glu Lys Gln Leu Gln Lys
20 25 30Asp Lys Gln Val Tyr Arg Ala Thr His Arg Leu Leu Leu Leu Gly Ala
35 40 45Gly Glu Ser Gly Lys Ser Thr Ile Val Lys Gln Met Arg Ile Leu His
50 55 60

Val

d'wnt.

65

<210> 84

<211> 58

<212> PRT

<213> Saccharomyces cerevisiae

<400> 84

Met Gly Cys Thr Val Ser Ala Glu Asp Lys Ala Ala Ala Glu Arg Ser
1 5 10 15

Lys Met Ile Asp Lys Asn Leu Arg Glu Asp Gly Glu Lys Ala Ala Arg
20 25 30

Glu Val Lys Leu Leu Leu Leu Gly Ala Gly Glu Ser Gly Lys Ser Thr
35 40 45

Ile Val Lys Gln Met Lys Ile Ile His Glu
50 55

<210> 85

<211> 58

<212> PRT

<213> Saccharomyces cerevisiae

<400> 85

Met Gly Cys Thr Val Ser Ala Glu Asp Lys Ala Ala Val Glu Arg Ser
1 5 10 15

Lys Met Ile Asp Arg Asn Leu Arg Glu Asp Gly Glu Lys Ala Ala Lys
20 25 30

Glu Val Lys Leu Leu Leu Leu Gly Ala Gly Glu Ser Gly Lys Ser Thr
35 40 45

Ile Val Lys Gln Met Lys Ile Ile His Glu
50 55

<210> 86

<211> 67

<212> PRT

<213> Saccharomyces cerevisiae

<400> 86

Met Ala Arg Ser Leu Thr Trp Arg Cys Cys Pro Trp Cys Leu Thr Glu
1 5 10 15

Asp Glu Lys Ala Ala Ala Arg Val Asp Gln Glu Ile Asn Arg Ile Leu
20 25 30

Leu Glu Gln Lys Lys Gln Asp Arg Gly Glu Leu Lys Leu Leu Leu Leu
35 40 45

Gly Pro Gly Glu Ser Gly Lys Ser Thr Phe Ile Lys Gln Met Arg Ile

cont.

50

55

60

Ile His Gly
65

<210> 87

<211> 66

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 87

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro
1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Arg Lys Leu Leu Leu Leu Gly
35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Ile Val Lys Gln Met Arg Ile Leu
50 55 60

His Val
65

<210> 88

<211> 66

<212> PRT

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<223> Description of Artificial Sequence: Synthetic
sequence

<400> 88

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro
1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Val Lys Leu Leu Leu Leu Gly
35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Ile Val Lys Gln Met Lys Ile Ile
50 55 60

His Glu
65

61
Gly

<210> 89
<211> 66
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
sequence

<400> 89
Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro
1 5 10 15
Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30
Leu Glu Lys Gln Arg Asp Lys Asn Glu Val Lys Leu Leu Leu Gly
35 40 45
Ala Gly Glu Ser Gly Lys Ser Thr Ile Val Lys Gln Met Lys Ile Ile
50 55 60
His Glu
65

<210> 90
<211> 66
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
sequence

<400> 90
Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro
1 5 10 15
Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30
Leu Glu Lys Gln Arg Asp Lys Asn Glu Leu Lys Leu Leu Leu Gly
35 40 45
Pro Gly Glu Ser Gly Lys Ser Thr Phe Ile Lys Gln Met Arg Ile Ile
50 55 60
His Gly
65

<210> 91
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 91

tgg cat tgg ttg cag cta aaa cct ggc cag cct atg tac
Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
1 5 10

39

<210> 92

<211> 13

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 92

Trp His Trp Leu Gln Leu Lys Pro Gly Gln Pro Met Tyr
1 5 10

<210> 93

<211> 39

<212> DNA

<213> *Saccharomyces cerevisiae*

<220>

<221> CDS

<222> (1)..(39)

<400> 93

tgg cat tgg ttg tcc ttg tgc ccc ggc cag cct atg tac
Trp His Trp Leu Ser Leu Ser Pro Gly Gln Pro Met Tyr
1 5 10

39

<210> 94

<211> 13

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 94

Trp His Trp Leu Ser Leu Ser Pro Gly Gln Pro Met Tyr
1 5 10

<210> 95

<211> 39

<212> DNA

<213> *Saccharomyces cerevisiae*

<220>

<221> CDS

<222> (1)..(39)

<400> 95

tgg cat tgg ttg tcc ctg gac gct ggc cag cct atg tac

39

Trp His Trp Leu Ser Leu Asp Ala Gly Gln Pro Met Tyr
1 5 10

<210> 96

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 96

Trp His Trp Leu Ser Leu Asp Ala Gly Gln Pro Met Tyr
1 5 10

<210> 97

<211> 39

<212> DNA

<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 97

ttg cat ttg ttg acc ttg atg gcc ggg cag cct atg tac
Trp His Trp Leu Thr Leu Met Ala Gly Gln Pro Met Tyr
1 5 10

39

<210> 98

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 98

Trp His Trp Leu Thr Leu Met Ala Gly Gln Pro Met Tyr
1 5 10

<210> 99

<211> 39

<212> DNA

<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 99

ttg cat ttg ttg cag ctg tcg gcg ggc cag cct atg tac
Trp His Trp Leu Gln Leu Ser Ala Gly Gln Pro Met Tyr
1 5 10

39

<210> 100

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 100

Trp His Trp Leu Gln Leu Ser Ala Gly Gln Pro Met Tyr
1 5 10

<210> 101

<211> 39

<212> DNA

<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 101

tgg cat tgg ttg agg ttg cag tcc ggc cag cct atg tac
Trp His Trp Leu Arg Leu Gln Ser Gly Gln Pro Met Tyr
1 5 10

39

<210> 102

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 102

Trp His Trp Leu Arg Leu Gln Ser Gly Gln Pro Met Tyr
1 5 10

<210> 103

<211> 39

<212> DNA

<213> Saccharomyces cerevisiae

<220>

<221> CDS

<222> (1)..(39)

<400> 103

tgg cat tgg ttg cgc ttg tcc gcc ggg cag cct atg tac
Trp His Trp Leu Arg Leu Ser Ala Gly Gln Pro Met Tyr
1 5 10

39

<210> 104

<211> 13

<212> PRT

<213> Saccharomyces cerevisiae

<400> 104

Trp His Trp Leu Arg Leu Ser Ala Gly Gln Pro Met Tyr
1 5 10

<210> 105
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>
<221> CDS
<222> (1)..(39)

<400> 105
tgg cat tgg ttg tgc ctc gtc ccg ggg cag cct atg tac
Trp His Trp Leu Ser Leu Val Pro Gly Gln Pro Met Tyr
1 5 10

39

<210> 106
<211> 13
<212> PRT
<213> Saccharomyces cerevisiae

<400> 106
Trp His Trp Leu Ser Leu Val Pro Gly Gln Pro Met Tyr
1 5 10

<210> 107
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>
<221> CDS
<222> (1)..(39)

<400> 107
tgg cat tgg ttg tcc ctg tac ccc ggg cag cct atg tac
Trp His Trp Leu Ser Leu Tyr Pro Gly Gln Pro Met Tyr
1 5 10

39

<210> 108
<211> 13
<212> PRT
<213> Saccharomyces cerevisiae

<400> 108
Trp His Trp Leu Ser Leu Tyr Pro Gly Gln Pro Met Tyr
1 5 10

<210> 109
<211> 39
<212> DNA
<213> Saccharomyces cerevisiae

<220>
<221> CDS

1
cont.

<222> (1)..(39)

<400> 109

tgg cat tgg ttg cgg cag cag ccc ggg cag cct atg tac
 Trp His Trp Leu Arg Leu Gln Pro Gly Gln Pro Met Tyr
 1 5 10

39

<210> 110

<211> 13

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 110

Trp His Trp Leu Arg Leu Gln Pro Gly Gln Pro Met Tyr
 1 5 10

<210> 111

<211> 62

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 111

Arg Ile Asp Thr Thr Gly Ile Thr Glu Thr Glu Phe Asn Ile Gly Ser
 1 5 10 15

Ser Lys Phe Lys Val Leu Asp Ala Gly Gly Gln Arg Ser Glu Arg Lys
 20 25 30

Lys Trp Ile His Cys Phe Glu Gly Ile Thr Ala Val Leu Phe Val Leu
 35 40 45

Ala Met Ser Glu Tyr Asp Gln Met Leu Phe Glu Asp Glu Arg
 50 55 60

<210> 112

<211> 62

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 112

Arg Val Leu Thr Ser Gly Ile Phe Glu Thr Lys Phe Gln Asn Asp Lys
 1 5 10 15

Val Asn Phe His Met Phe Asp Val Gly Gly Gln Arg Asp Glu Arg Lys
 20 25 30

Lys Trp Ile Gln Cys Phe Asn Asp Val Thr Ala Ile Ile Phe Val Val
 35 40 45

Ala Ser Ser Ser Tyr Asn Met Val Ile Arg Glu Asp Asn Gln
 50 55 60

<210> 113

<211> 62
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 113
Arg Val Lys Thr Thr Gly Ile Val Glu Thr His Phe Thr Phe Lys Asp
1 5 10 15
Leu His Phe Lys Met Phe Asp Val Gly Gly Gln Arg Ser Glu Arg Lys
20 25 30
Lys Trp Ile His Cys Phe Glu Gly Val Thr Ala Ile Ile Phe Cys Val
35 40 45
Ala Leu Ser Ala Tyr Asp Leu Val Leu Ala Asp Glu Glu Met
50 55 60

<210> 114
<211> 62
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 114
Arg Val Lys Thr Thr Gly Ile Val Glu Thr His Phe Thr Phe Lys Asp
1 5 10 15
Leu Tyr Phe Lys Met Phe Asp Val Gly Gly Gln Arg Ser Glu Arg Lys
20 25 30
Lys Trp Ile His Cys Phe Glu Gly Val Thr Ala Ile Ile Phe Cys Val
35 40 45
Ala Leu Ser Asp Tyr Asp Leu Val Leu Ala Glu Asp Glu Glu
50 55 60

<210> 115
<211> 62
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 115
Arg Val Lys Thr Thr Gly Ile Val Glu Thr His Phe Thr Phe Lys Asn
1 5 10 15
Leu His Phe Arg Leu Phe Asp Val Gly Gly Gln Arg Ser Glu Arg Lys
20 25 30
Lys Trp Ile His Cys Phe Glu Asp Val Thr Ala Ile Ile Phe Cys Asn
35 40 45
Ala Leu Ser Gly Tyr Asp Gln Val Leu His Glu Asp Glu Thr
50 55 60

<210> 116

<211> 62
<212> PRT
<213> Saccharomyces cerevisiae

<400> 116

Arg Val Pro Thr Thr Gly Ile Ile Glu Tyr Pro Phe Asp Leu Glu Asn
1 5 10 15

Ile Ile Phe Lys Met Val Asp Ala Gly Gly Gln Arg Ser Glu Arg Lys
20 25 30

Lys Trp Ile His Cys Phe Glu Asn Val Thr Ser Ile Met Phe Leu Val
35 40 45

Ala Leu Ser Glu Tyr Asp Gln Cys Leu Glu Glu Asn Asn Gln
50 55 60

<210> 117

<211> 62

<212> PRT

<213> Saccharomyces cerevisiae

<400> 117

Arg Met Pro Thr Thr Gly Ile Asn Glu Tyr Cys Phe Ser Val Gln Lys
1 5 10 15

Thr Asn Leu Lys Ile Val Asp Ala Gly Gly Gln Arg Ser Glu Arg Lys
20 25 30

Lys Trp Ile His Cys Phe Glu Asn Ile Ile Ala Leu Ile Tyr Leu Ala
35 40 45

Ser Leu Ser Glu Tyr Asp Gln Val Leu Val Glu Ser Asp Asn
50 55 60

<210> 118

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> The 5'-end of this sequence is linked to the
3'-end of SEQ ID NO: 21 by (NNN) n where n is any
chosen integer

<220>

<223> Description of Artificial Sequence: Synthetic
sequence

<400> 118

agcttctgcc tcacgcttaa gtagc

25

<210> 119

<211> 26

<212> DNA
<213> Artificial Sequence

<220>
<223> The 5'-end of this sequence is linked to the
3'-end of SEQ ID NO: 25 by (NNN) n where n is any
chosen integer

<220>
<223> Description of Artificial Sequence: Synthetic
sequence

<400> 119
gttgtccttc ttttcactcg agtacc

26

<210> 120
<211> 9
<212> PRT
<213> Saccharomyces cerevisiae

<400> 120
Leu Leu Leu Leu Gly Ala Gly Glu Ser
1 5

<210> 121
<211> 9
<212> PRT
<213> Saccharomyces cerevisiae

<400> 121
Leu Glu Lys Gln Arg Asp Lys Asn Glu
1 5

<210> 122
<211> 6
<212> PRT
<213> Saccharomyces cerevisiae

<220>
<221> VARIANT
<222> 2,4,5
<223> Xaa may be any amino acid

<400> 122
Gly Xaa Gly Xaa Xaa Gly
1 5

<210> 123
<211> 10
<212> PRT
<213> Saccharomyces cerevisiae

<400> 123
Leu Leu Leu Leu Gly Ala Gly Glu Ser Gly

21
cont.

1

5

10

<210> 124

<211> 6

<212> PRT

<213> Saccharomyces cerevisiae

<220>

<221> VARIANT

<222> (3)...(6)

<223> Xaa may be any amino acid

<400> 124

Met Gly Xaa Xaa Xaa Ser

1

5

<210> 125

<211> 9

<212> PRT

<213> Saccharomyces cerevisiae

<400> 125

Gly Ser Gly Glu Ser Gly Asp Ser Thr

1

5

<210> 126

<211> 8

<212> PRT

<213> Saccharomyces cerevisiae

<400> 126

Gln Ala Arg Lys Leu Gly Ile Gln

1

5

<210> 127

<211> 9

<212> PRT

<213> Saccharomyces cerevisiae

<400> 127

Leu Ile His Glu Asp Ile Ala Lys Ala

1

5

<210> 128

<211> 7

<212> DNA

<213> Saccharomyces cerevisiae

<400> 128

tgaaaca

<210> 129

<211> 10
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 129
Leu Leu Leu Leu Gly Ala Gly Glu Ser Gly
1 5 10

<210> 130
<211> 8
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 130
Leu Leu Leu Leu Gly Ala Gly Glu
1 5

<210> 131
<211> 6
<212> DNA
<213> *Saccharomyces cerevisiae*

<400> 131
gaggct

6

<210> 132
<211> 4
<212> DNA
<213> *Saccharomyces cerevisiae*

<400> 132
gaga

4

<210> 133
<211> 11
<212> PRT
<213> *Saccharomyces cerevisiae*

<400> 133
Tyr Ile Ile Lys Gly Val Phe Trp Asp Pro Ala
1 5 10

D
word.

E